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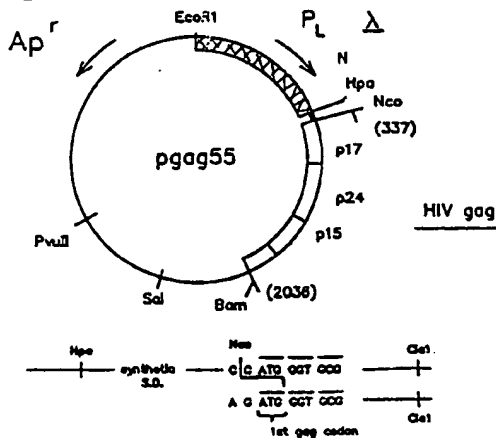
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⑤④ **Human immunodeficiency virus GAG-encoded proteins.**

⑤⑦ The invention concerns nonfused recombinant proteins corresponding to the complete HIV gag and the complete HIV p17 subregion coding sequences, which are useful immunological components of diagnostics, therapeutics, and vaccines.

FIG. 2



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Human Immunodeficiency Virus gag-encoded Proteins

FIELD OF THE INVENTION

The present invention concerns proteins derived by molecular cloning that are useful immunological components of diagnostics, therapeutics, and vaccines for infectious diseases. In particular, the invention relates to proteins encoded by the gag gene of human immunodeficiency virus, the etiologic agent of acquired immunodeficiency syndrome.

BACKGROUND

The human immunodeficiency virus (HIV, also LAV, HTLV-III, or ARV) is the primary etiologic agent of the acquired immune deficiency syndrome (AIDS) [Barre-Sinoussi et al., *Science* 220:868-871 (1983); Gallo et al., *Science* 224:500-503 (1984); Levy et al., *Science* 225:840-842 (1984)]. The underlying disease state involves a tropism of HIV for the T4 lymphocyte subset resulting in a selective depletion of the helper/inducer cells of the immune system, leaving the HIV-infected individual defenseless against a number of opportunistic infections. Approximately two million people in the United States and five million or more individuals world-wide may now be infected by HIV. The U.S. Public Health Service now estimates that the cost of caring for AIDS patients in 1991 will be between \$8 billion to \$16 billion per year. Thus, the development of diagnostics, therapeutics, and vaccines to HIV is the subject of intense medical research.

The nucleotide sequence of several independent viral isolates of HIV have been determined [Ratner et al., *Nature* 313:277-284 (1985); Muesing et al., *Nature* 313:250-258 (1985); Wain-Hobson et al., *Cell* 40:9-17 (1985); Sanchez-Pescador et al., *Science* 227:484-492 (1985)]. HIV is related to the lentivirus subgroup of retroviruses. Lentiviruses are associated with slowly developing, non-tumorigenic diseases in other animal systems. The major structural proteins of the HIV infectious particle are encoded by the viral gag, pol, and env genes. In addition to the gag, pol, and env genes, the genome of HIV contains several other open reading frames designated src, tat, art, tr, and tr which encode additional viral proteins. These proteins are known to serve important regulatory functions during the HIV infectious cycle.

The gag gene product are involved in the

structure of the genomic RNA-containing core and the assembly of the virus. The gag gene is termed "gag" for group-specific antigen, since in other retroviral systems the analogous gene encodes proteins that are cross-reactive with similar proteins of a related group of retroviruses. The HIV gag gene encodes a precursor of about 55,000 daltons (55 kDa) which is designated p55^{gag}. The gag coding sequence of HIV isolate BH10 (HIV_{BH10}) (Ratner et al., *op. cit.*) is contained within nucleotides 334 to 1869 and is comprised of 512 codons (Figure 1). The nucleotide numbering is according to Ratner et al. (*op. cit.*).

During the HIV infectious life cycle, the gag precursor p55^{gag} is proteolytically processed into species of approximately 24, 17, and 15 kDa. The gag proteins are designated p24^{gag}, p17^{gag}, and p15^{gag}. The mature gag proteins are derived from subregions of the p55^{gag} precursor in the order N-p17-p24-p15-C (N- and -C designate the amino and carboxyl terminus of the protein, respectively).

As is the case for the N-terminal gag-encoded proteins of other mammalian retroviruses [Henderson et al., *Proc. Natl. Acad. Sci.* 80:339-343 (1983); Rein et al., *Proc. Natl. Acad. Sci.* 83:7246-7250 (1986)], the N-terminal gag-encoded p17^{gag} is modified at its mature N-terminal glycine by myristylation. The N-terminal lipid myristic acid moiety of p17^{gag} presumably mediates the interaction of p17^{gag} with the lipid membrane of the virus particle. Gag-derived determinants are known to be exposed on the surface of cells infected by other mammalian retroviruses [Schiff-Maker and Rosenberg, *Virology* 154:286-301 (1986)]. Thus, HIV p17^{gag}-specific epitopes may be present on the surface of infectious HIV particles and HIV-infected cells. The p24^{gag} protein is the major virus capsid or core protein. The C-terminal gag-encoded p15^{gag} protein is highly basic and likely constitutes the core ribonucleoprotein and binds nonspecifically to many sites on the viral RNA.

Earlier reports describe the expression of subregions of HIV gag protein in heterologous expression systems such as in *E. coli* [Dowbenko et al., *Proc. Natl. Acad. Sci.* 82: 7748-7752 (1985); Chang et al., *Science* 228:93-99 (1985); Ghayeb et al., *DNA* 5:93-99 (1986); Steimer et al., *Virology* 150: 283-290 (1986); Shoeman et al., *Anal. Biochem.* 161:370-379 (1987)] and in yeast [Kramer et al., *Science* 231:1580-1584 (1986)]. These recombinant gag-derived proteins produced in heterologous expression systems were shown to be useful as antigens for the detection of HIV gag-specific antibodies in human sera. No previous report has described the production using a het-

erologous expression system, for example an *E. coli* plasmid expression system, of proteins corresponding to the full length, nonfused gag precursor p55^{gag} or to the mature gag subregion p17^{gag}. The present invention provides the full length nonfused HIV gag precursor p55^{gag} and the mature gag subregion p17^{gag} as produced using a heterologous expression system.

SUMMARY OF THE INVENTION

The invention provides proteins corresponding to the full length, nonfused HIV gag precursor p55^{gag} and the mature nonfused p17^{gag} subregion of p55^{gag} which are produced using molecular cloning methods and heterologous expression vector systems. Polypeptides corresponding to p55^{gag} and p17^{gag} of the HIV isolate BH10 (Ratner et al., *op. cit.*) were expressed in *E. coli*. Protein p55^{gag} is encoded by nucleotides number 334 to 1869 (nucleotide numbering according to Ratner et al., *op. cit.*) (see Figure 1). The p17^{gag} segment of p55^{gag} is encoded by nucleotides number 334 to 729. The HIV gag proteins are useful immunological components of diagnostics to identify individuals and blood products that have been exposed to HIV, of reagents that can be used to monitor and stage the progress of the disease in clinical therapeutic trials, of therapeutic agents able to control the disease and of a vaccine that will protect individuals who are exposed to HIV.

DETAILED DESCRIPTION OF THE INVENTION

Two recombinant peptides containing only amino acid sequences corresponding to the gag coding sequence of HIV have been created. These recombinant gag-derived polypeptides are highly immunoreactive with human sera from individuals infected with HIV. The term "peptide" is well known in the art and refers to a compound of two or more amino acids. One of the peptides of the invention corresponds to the complete and nonfused HIV gag precursor protein. In HIV_{BH10}, p55^{gag} is encoded by nucleotides 334 to 1869 (nucleotide numbering according to Ratner et al., *op. cit.*). A second peptide of the invention corresponds to the mature 4-terminal gag-encoded protein, p17^{gag}. In HIV_{BH10}, p17^{gag} is encoded by nucleotides 334 to 729 (Ratner et al., *op. cit.*). The specific domains of HIV are known in the art.

The present *E. coli* expressed proteins corresponding to HIV p55^{gag} and p17^{gag} are referred

to as GAG55 and GAG17, respectively. The structures of plasmid expression vectors designed for the expression of GAG55 and GAG17 in *E. coli* are detailed in Example 1.

As used herein, the term "expression vector" includes a DNA molecule which contains signals, recognized by a particular host biological cell, that direct and determine the expression of a particular gene sequence in the host cell. The vector DNA may, for example, contain information that determines the uptake of the vector DNA by the host cell, the integration and replication of the vector DNA, and the corresponding RNA biosynthesis and translation. Examples of expression vectors include recombinant viruses, plasmids and genes.

It is to be understood that the term "corresponding to" includes modifications of the specified amino acid sequences which do not adversely affect the antigenic characteristics of the peptide of the invention. For example, different isolates of HIV are known to differ to some extent in the predicted amino acid sequence of the gag-derived proteins (Ratner et al., *op. cit.*; Muesing et al., *op. cit.*; Wain-Hobson et al., *op. cit.*; Sanchez-Pescador, *op. cit.*). One skilled in the art could align the amino acid sequences of peptides from different sources to the schematic of Figure 1 or the nucleotide sequence of Ratner et al. (*op. cit.*) to identify the segments therein which correspond to the peptides defined herein.

HIV gag-derived proteins are useful immunological components of diagnostics, therapeutics, and vaccines for HIV-related disease, including acquired immune deficiency syndrome (AIDS). HIV-expressed gag proteins are known to be useful immunological antigens for the detection of HIV gag-specific antibodies [Petricciani, *Ann. Int. Med.* **103**:726-729 (1985)]. The analysis of HIV-specific antibodies in human serum samples is extremely important in diagnosis of disease and for screening of human blood products. An enzyme-linked immunosorbent assay (ELISA) which uses HIV grown in tissue culture cells as antigen is now widely used for detecting sera positively immunoreactive to HIV (Petricciani, *op. cit.*). The ELISA derived from purified HIV particles grown in tissue culture is designated HIV-ELISA.

Since HIV infection leads to a series of clinical manifestations accompanied by variable antibody levels to different antigens, useful blood screening and diagnostic reagents should contain multiple antigens. Moreover, it is desirable to detect and quantitate antibody levels to specific antigenic components of HIV. Although the HIV-ELISA is a sensitive tool for the detection of HIV-specific antibodies in human serum samples, this test does not provide information as to which antigenic components of HIV are reactive with human antibodies.

The production of individual defined HIV-derived proteins, such as gag-encoded proteins, in E. coli provides antigen reagents for the selective detection and quantitation of antibodies recognizing specific antigens, such as gag-encoded antigens. HIV gag proteins derived by recombinant cloning and expressed in heterologous expression systems, i.e., recombinant gag proteins, are suitable for the development of novel HIV diagnostic kits. For example, the E. coli produced HIV GAG55 and GAG17 proteins can be efficiently used in a diagnostic ELISA for the detection and quantitation of HIV gag-specific antibodies that are associated with prior infection with HIV.

A heterologous expression system such as that utilizing an E. coli plasmid expression vector, provides a preferred alternative to infectious HIV grown in tissue culture as a source of antigen for ELISA blood screening and diagnostic reagents. Importantly, the E. coli plasmid expression system is free of infectious HIV. Moreover, gag-derived proteins GAG55 and GAG17 can be expressed at high levels in E. coli using the appropriate vectors, thereby facilitating the purification of these useful polypeptides. Thus, recombinant gag proteins can be produced in large quantities without the biohazard and containment problems associated with the production of gag proteins from HIV virions.

The recombinant gag proteins of the present invention are preferred to the previously described recombinant gag proteins. The recombinant product GAG55 corresponds to the complete, non-fused product of the complete gag coding sequence. Previously described recombinant gag proteins only contain subregions of the gag coding sequence. For example, in a recombinant gag protein described by Kramer et al. (op. cit.) and Shoemaker et al. (op. cit.), amino acids number 2 to 12 of the gag coding sequence were deleted. Other reported recombinant gag proteins have reported have larger segments of the gag coding sequence deleted (Dowbenko et al., op. cit.; Ratner et al., op. cit.; Ghrayeb et al., op. cit.; Steinhilber et al., op. cit.). More gag-specific antigenic determinants are represented by the product of the complete full length gag coding sequence than are represented by incomplete subregions of gag. The product of the full length gag coding sequence, such as GAG55, is expected to be a more sensitive reagent for the detection of gag-specific antibodies than are antigens comprised of only subregions of gag.

In particular, GAG55 and GAG17 contain the region of gag corresponding to the complete p17^{gag}. This complete region is not present in other reported recombinant gag proteins. As discussed previously and shown in the p17^{gag} region of gag contains antigenic determinants that may be presented on the surface of HIV-infected cells, and may be critical for a protective host

immune response to HIV.

Another advantage of the peptides of the invention is that the recombinant gag-derived proteins are not fused to other, non-HIV, or non-HIV gag sequences. The inclusion of additional non-HIV sequences, such as E. coli-derived sequences, fused to the HIV gag coding sequence is not desirable since these sequences could lead to frequent false-positive (i.e., non-HIV) antibody reactivity. Thus, the HIV gag sequences have been expressed in E. coli in the absence of fused bacterial or other non-HIV protein sequences. It is also desirable to express nonfused authentic gag proteins, rather than fusion proteins, in order to facilitate the correct folding and native structure of the product. The correctly folded structure of the product may be critical to its effectiveness as a diagnostic, therapeutic, or vaccine reagent. Previously reported recombinant HIV gag-derived proteins are fusion polypeptides containing non-HIV sequences.

Although most current strategies for the development of an AIDS vaccine are focusing on the HIV env gene, the gag antigens may well be an important component of a protective vaccine against HIV. Indeed, recent reports suggest that gag p17^{gag} antigenic determinants are accessible to the neutralizing effects of specific antibodies [Sarin et al., *Science* 232: 1135-1137 (1988)]. As discussed previously, p17^{gag} determinants may be exposed on the envelope of the virus itself, since the gag p17^{gag} protein is thought to be associated with the virus lipid membrane.

AIDS patients are less likely to have antibodies to gag-encoded proteins compared to individuals with milder forms of the disease [Steimer et al., *Virology* 150: 283-290 (1986)]. Thus, there is at least correlative evidence that the gag-specific host immune response may be critical for the effective control of HIV by an HIV-exposed individual.

Due to the heterogeneity in the sequence of env from various HIV isolates [Ratner et al., *Nature* 313:277-284 (1985); Wain-Hobson et al., *Cell* 40:9-17 (1985); Muesing et al., *Nature* 313:250-258 (1985); Sanchez-Pescador et al., *Science* 227:484-492 (1985)], it is possible that a vaccine containing env from a single HIV isolate may not have a broadly neutralizing effect on other HIV species. The degree of heterogeneity in the gag gene product among different HIV isolates is far less than that found for env [Ratner et al., *Nature* 313:277-284 (1985); Muesing et al., *Nature* 313:250-258 (1985); Wain-Hobson et al., *Cell* 40:9-17 (1985); Sanchez-Pescador et al., *Science* 227:484-492 (1985)]. Thus, HIV gag gene products may be an important component of a vaccine that is broadly protective against many isolates of HIV.

Since the peptides of the invention are produced by recombinant DNA technology in the ab-

sence of infectious HIV, they provide several advantages over more traditional vaccine approaches using killed or attenuated virus preparations. Recombinant peptides are safe to prepare and administer and no genetic information is introduced by vaccination. In addition recombinant peptides are more readily produced and purified than the viral-expressed peptides isolated from virus-infected cells.

In one embodiment, the peptide of the invention is used in a diagnostic kit used for detecting antibodies to HIV in a biological sample. The peptide can be employed in a process for detecting antibodies to HIV comprising contacting a biological sample with the peptide and detecting immunoreactivity. The peptide can also be employed as a component in a vaccine protective against HIV. The vaccine comprises an effectively protective amount of the peptide.

E. coli strain KA1298, transformed with plasmid pGAG55, encoding protein C, is described in Example 1, has been deposited with the American Type Culture Collection (ATCC), Rockville, Maryland, and bears deposit accession number 67378. This deposit is available to the public upon the grant of a patent to the assignee. However, it should be understood that the availability of a deposit does not constitute an offer to practice the subject matter in derogation of patent rights granted by governmental authority.

Materials and Methods

Unless otherwise specified, parts and percentages are by weight and temperature are Celsius.

Plasmid Construction

Plasmid construction were carried out using standard methodology as described by Maniatis et al., Molecular Cloning, 2nd ed., by Plasmid, Cold Spring Harbor Laboratory Press, New York (1982), the teaching of which is hereby incorporated by reference. Enzymes and other reagents used for plasmid constructions were obtained from Bethesda Research Laboratories, Gaithersburg, MD or New England Biolabs, Beverly, MA. Methods for digesting, identifying, recovering and purifying the various nucleotide sequences set forth in the invention are known to those skilled in the art as are methods for ligating the various vectors, transforming host microorganisms, cloning, and recovering products. Accordingly, the methods will only be described by reference to specific embodiments of the invention set forth hereinafter.

E. coli Strains, Plasmid Expression Vectors, and Induction of Expression

Several types of *E. coli* plasmid expression vectors were used for the production of HIV gag-derived proteins in *E. coli*. In the vectors used, expression of the desired product is controlled transcriptionally using either the operator and promoter of the *E. coli* tryptophan operon [Ivanoff et al., Proc. Natl. Acad. Sci. 83:5392-5396 (1986)] or the phage lambda P_L promoter [Rosenberg et al., Methods Enzymol. 101:123-138 (1983)]. We have also used the *E. coli* lactose operon promoter to direct expression of HIV gag-derived proteins in *E. coli*.

Plasmid vectors utilizing the phage lambda promoter were derived from plasmid pBF106. The *E. coli* plasmid expression vector pBF106 was constructed as follows. The small HpaI to BamHI fragment of plasmid pKC30 [Rosenberg et al., Methods Enzymol. 101:123-138 (1983); Rao, Gene 31:247-150 (1984)] (obtained from K. Abremski, Du Pont Experimental Station, Wilmington, DE) was replaced with the following oligonucleotide linker:

5'- AAC GAA TCC GAA GTG TAA GCC ATG -3'
3'- TTG CTT AGG CTT CAC ATT CGG TAC CTAG -5'

The inserted sequence provides signals for efficient translation initiation and a unique NcoI restriction endonuclease site. Since the translation start ATG forms part of the NcoI site, coding sequences can easily be fused in-frame directly to the ATG and translation initiation signal.

Expression from the P_L-containing plasmids is controlled transcriptionally using a lysogenic host that provides lambda cI repressor. Expression is induced by inactivating cI using nalidixic acid [Mott et al., Proc. Natl. Acad. Sci. 82:88-92 (1985)] or by temperature-shift and using a temperature sensitive repressor such as cI857 [Rosenberg et al., Methods Enzymol. 101:123-138 (1983)]. Temperature-shift inductions were carried out using host strain KA1298 (obtained from K. Abremski, Du Pont Experimental Station, Wilmington, DE), according to the procedure of Young et al., Proc. Natl. Acad. Sci., 80:6105-6109 (1983). KA1298 is a defective lambda lysogenic containing cI857. Nalidixic acid inductions were carried out using the defective lysogenic host DC550 (obtained from D. Court, NCI, Frederick, MD), according to Mott et al., Proc. Natl. Acad. Sci. 82:88-92 (1985).

E. coli plasmid expression vectors containing the tryptophan promoter were derived from pKGP36.trp (Ivanoff et al., op. cit.). *E. coli* host strains MM294 and HB101, DH5 (F⁻ recA1, endA1, gyrA96, hsdR17, supE44), DH5Δ (lac), SC122, CAG456(htrR165), CAG629(lon-htrR165), and SG4119 (Δlon) were used [Maniatis et al., Molecu-

lar Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, NY (1982); [Ivanoff et al., *Proc. Natl. Acad. Sci.* 81:6779-6783]. The expression of the trypanophan promoter was carried out as described (Ivanoff et al., *op. cit.*).

Serum Samples and Serology

Human sera were collected at Food Bank of Delaware blood donor centers between 1986 or were obtained from T. Mathews, Duke University. Serum was stored at -70°C before testing and repeated freezing and thawing was avoided. Preliminary classification of all sera was made based on repeated tests with an HIV-ELISA (Du Pont, Wilmington, DE) blood screening kit, on the results of HIV immunoblot testing carried out by commercial laboratory (Biotech Research Laboratories, Rockville, MD), and on testing sera for reactivity to env using a recombinant antigen (EM) based ELISA (described below). Procedure for 3-1/2 hour assay described in the Du Pont HIV-III ELISA blood screening kit, was used to test sera. Further immunoblot analysis was carried out using commercially available HIV anti- β -containing nitrocellulose strips (Du Pont-Biotec Research Laboratories).

Immunoblot Analysis

Standard methods for polyacrylamide gel electrophoresis (SDS-PAGE) [Laemmli, *Biochem. J.* 227:680-685 (1970)] and electrophoretic transfer of proteins to nitrocellulose [Towbin et al., *Proc. Natl. Acad. Sci.* 76:4350-4354 (1979)] were used. Gel preparations of GAG55 (30 ng of total protein per lane) were typically used for immunoblot analysis. Following electrophoretic transfer, nitrocellulose strips were preincubated 30 minutes at room temperature in Blotto buffer (1% w/v non-fat milk solids, 0.05% Tween 20, 0.1% v/v goat serum, phosphate buffered saline, pH 7.4). Strips were then incubated with a dilution of sample (1:50 in Blotto buffer) or sample control (1:50 in Blotto buffer) overnight at 4°C . Strips were then washed exhaustively with phosphate buffered saline containing 0.05% Tween 20. Human immunoglobulins were detected using biotinylated anti-human IgG (Vector Laboratories, Burlingame, CA) and avidinD-horseradish peroxidase (Vector Laboratories, Burlingame, CA). Competition of sera for competition experiments was carried out by incubation of 10 to 20 μl of serum with 100 μl of *E. coli* extract (containing 100 μl of bacterial protein) for 2 hours at 25°C followed by centrifugation to remove insoluble material.

Protein Purification

E. coli-expressed GAG55 was purified by immunoaffinity chromatography. Induced *E. coli* were collected by centrifugation and the cell pellet was stored at -70°C . The induced *E. coli* cell pellet was suspended in PBS and the cells were lysed using a French press. The lysate was centrifuged (27000 x g) and the supernatant applied to an immunoaffinity column containing p17^{gag} mouse monoclonal-specific antibody BT2 (Biotech Research Laboratory, Rockville, MD) covalently attached to Reacti-gel (Pierce Chemical, Rockford, IL). The immunoaffinity column was prepared as specified by the supplier and washed and equilibrated with phosphate buffered saline (PBS). The *E. coli* cell lysate was loaded onto the column and the column was extensively washed with PBS. Fractions containing gag protein were eluted from the column using 0.5 N acetic acid and were then pH neutralized, lyophilized, and dissolved in PBS containing 0.1% SDS.

Alternatively, GAG55 was purified from a French press lysate of *E. coli* by cation exchange chromatography. Soluble proteins were dialyzed against 0.05 M phosphate buffer (pH 7.0), applied to a Pharmacia Mono S column (Pharmacia, Piscataway, NJ) and eluted with a NaCl gradient. The purity of the antigen was approximately 90% purified.

The peptide ENV9 was purified from *E. coli* containing the vector pENV9, as discussed in assignee's copending application, serial number 010056, filed Feb. 2, 1987. The predicted protein product of pENV9, designated ENV9, contains 340 amino acids which correspond to 54 amino acids from the N-terminal of the poliovirus sequence of pEXC [Ivanoff et al., *op. cit.*], 46 amino acids of the C-terminal of the HIV env gp120 domain and 340 amino acids of the N-terminal of the env gp41 domain. Enzyme Linked Immunosorbent Assay (ELISA)

Purified GAG55 antigen in 60 mM Na_2CO_3 pH 9.6 buffer containing 0.0005% SDS was applied to Immulon II microtiter plates at 100 ng/well at 4°C for about 18 hours. Purified ENV9 antigen was similarly applied in 60 mM carbonate pH 9.6 buffer containing 0.00006% SDS at a concentration of 20 ng/well. ELISA procedures were carried out as specified for the commercial HIV-ELISA kit (Du Pont, Wilmington, DE). The antigen-coated plates were washed with PBS + 0.05% Tween 20 (PBS-T). Plate washings were performed using 3 cycles on a Titertek Microplate Washer 120 followed by rotating the plate and washing again. The plates were incubated with PBS-T for 1 hour at 37°C and were then washed 3 times with PBS-T and stored dry at 4°C until they were used.

The plates were incubated with patient sera at

a 1:20 to 1:100 dilution in 100 μ l of bovine serum albumin (1% w/v), 0.5% normal goat serum (NGS, 1% w/v), 0.05% thimerosal) in the microtiter wells. The plates were incubated for 2 hours at 25°. The plates were washed with PBS-T, exposed to goat anti-human IgG conjugated to alkaline phosphatase (Miles, ImmunoResearch, West Chester, OH), and washed with PBS-T. The color reaction was developed by the supplier, by exposure to 10% 4-chloro-1-naphthol phosphate in 100 μ l of 0.1 M Tris buffer (2 M, pH 9.8) with magnesium chloride (0.02% sodium azide) at a for 3 minutes, followed by addition of sodium hydroxide (1 N). The plates were read by a Multiskan MCC microtiter plate reader at 405 nm.

BRIEF DESCRIPTION OF FIGURES

FIGURE 1: Nucleotide sequence of the *gag* region of HIV_{BR10} (Rattus norvegicus). The start of the *gag* p55^{gag} coding sequence is indicated by an open bracket and the end of the *gag* p55^{gag} coding sequence is indicated by a closed bracket. The junctions between the p17^{gag} and p24^{gag} polypeptides, the site of the proteolytic processing of p55^{gag}, is indicated by a vertical line. The p55^{gag} is comprised by *gag* amino acid residues 1 to 512; p17^{gag} is comprised by amino acid residues 1 to 132.

FIGURE 2: Schematic representation of *E. coli* plasmid vector derived from the full length, non-fused HIV *gag* precursor (pGAG55). Plasmid pGAG55 (pgag55) is derived from pBR322 (thin line), pGAG55 (thick cross-hatched line), and pGAG55 (thick line), as well as a chemically synthesized segment (thick closed line). In the *gag* coding sequence, the phage lambda P_L promoter and the phage lambda P_R promoter are indicated. The *gag* coding sequence is preceded by a unique NcoI site. The *gag* ATG start codon has been changed to a unique NcoI site.

FIGURE 3: Expression of *E. coli* pGAG55. The accumulation of pGAG55 was analyzed by SDS-PAGE and Coomassie staining. Lane 1: DC550/pBF106 (DC550/pBF106) containing pBF106 (DC550/pBF106) in the presence of nalidixic acid. Lane 2: DC550/pGAG55 (DC550/pGAG55) in the presence of nalidixic acid. Lane 3: DC550/pGAG55, in the presence of nalidixic acid. The

GAG55 protein represents approximately 5% of total cell protein in the nalidixic acid-induced DC550 cells containing pGAG55 (lane 3).

FIGURE 4: Immunoblot reactivity of *E. coli*-expressed GAG55-derived proteins with human AIDS patient serum antibodies. Total *E. coli* lysates were analyzed. Lane 1: DC550/pGAG55, non-induced; lane 2: DC550/pGAG55 nalidixic acid-induced; lane 3: DC550/pBF106, nalidixic acid-induced.

FIGURE 5: Binding of *gag*-specific antibodies in human HIV-seropositive sera by *E. coli*-expressed GAG55, as analyzed by competition binding and immunoblot. Prior to immunoblot of HIV antigens (Blotech Research Labs, Rockville, MD), human HIV-seropositive sera were preincubated with cell extracts prepared from induced *E. coli*, as indicated. Lane 9: DC550/pBF106, 20 μ l extract; lane 10: DC550/pGAG55, 5 μ l extract; lane 11: DC550/pGAG55, 20 μ l extract. As shown, *E. coli*-expressed GAG55 protein effectively and selectively binds to all of the immunoblot-detectable *gag*-specific antibodies in this AIDS patient serum. Similar results have been obtained for several other HIV-seropositive sera. Thus, essentially all of the HIV *gag*-derived determinants and epitopes recognized by human antibodies are presented by *E. coli*-expressed GAG55. The figure also illustrates the identification of human serum samples containing HIV *gag*-reactive, HIV *env*- and *pol*-nonreactive antibodies. Human sera were analyzed by immunoblot of HIV antigens. In lanes 2 to 8, the specificity of the immunoblot reactivity for *gag* antigen was confirmed by competition with *E. coli*-expressed GAG55 protein (not shown).

FIGURE 6: Frequency distribution of GAG55-ELISA reactivity in three sample populations. Histograms show the fractions of samples in the absorbance ranges indicated on the horizontal axis. Absorbance ratio is ratio of the sample absorbance to that of a reference HIV positive control serum. Samples with absorbance ratios greater than 0.5 are grouped together for clarity. Upper panel: 69 normal HIV-ELISA-nonreactive blood donors. Middle panel: 57 representative HIV-ELISA reactive *env*-nonreactive sera from a blood donor population. Lower panel: 31 HIV positive sera: 26/31 of these sera are from patients with clinically diagnosed AIDS or AIDS related complex. Statistics for the populations were: normal sera, mean absorbance ratio = 0.00, s.d. = 0.03; HIV-reactive blood donors, mean = 0.13, s.d. = 0.11; HIV positives, mean = 0.37, s.d. = 0.43.

FIGURE 7: Comparison of apparent HIV *gag*-specific and *env*-specific antibody levels in human sera analyzed by ELISA. Antigens ENV9 and GAG55 were purified following expression in *E. coli*. Symbols: open triangles, HIV-seronegative; open

circles, non-AIDS, HIV-seropositive, and AIDS patients HIV-seropositive. Apparent gag-specific antibody levels are significantly higher in AIDS patient sera compared to non-AIDS HIV-seropositive sera. This apparent increase in gag-specific antibody levels is due to the fact that env-specific antibody levels are not significantly affected.

EXAMPLE

The invention is further described in the following Examples, wherein parts are by weight and degrees are by weight and degrees are by weight.

EXAMPLE

Plasmid Constructions

HIV sequences were obtained from clone λ -BH10 (Ratner et al., 1982) containing SacI fragment of HIV DNA isolated and treated with DNAase. The limited digestion to generate fragments. The randomly cut HIV DNA fragments were blunt ended by Klenow reaction, ligated with linker, cut with HindIII, and inserted into HindIII site of plasmid pTORF2. pTORF2 is derived from pBR322 (BamHI to NcoI fragment) containing HIV-derived sequences inserted at the transcription start site, tryptophan operator and promoter (see, e.g., op. cit.). The resulting plasmids were used to transform *E. coli* cells. The transformants containing HIV-derived sequences were identified by colony hybridization using the HIV_{BH10} proviral DNA as a probe. The plasmids containing HIV-derived sequences were designated pGAG3, pGAG4, PR2. pGAG3 and pGAG4 were shown by DNA sequencing to contain HIV-derived DNA inserted from nucleotides 227 to 1010, 227 to 2037, respectively. HIV nucleotide numbering according to Ratner et al. (1982). The gag coding sequence of pGAG3 contains nucleotides 334 to 1869 (see, e.g., Ratner et al., 1982) and pGAG4 contains the 5' end of the gag coding sequence and the HIV-derived sequence beyond the 3' end of the gag coding sequence.

Plasmids pGAG9 and pGAG10 specify the expression in *E. coli* under the control of the

control of the complete HIV_{BH10} gag coding sequence, were derived as described below. pGAG9 was derived by ligation of the following DNA fragments: 1) the PstI to PstI tryptophan promoter-containing fragment of pGAG4, 2) the PstI to PstI pBR322 origin-containing fragment from pR2. pGAG10 was similarly derived by ligation of the following DNA fragments: 1) the PstI to PstI tryptophan promoter-containing fragment of pGAG3, 2) the PstI to PstI pBR322 origin-containing fragment of pR2. In this way the PstI site within the gag coding sequence was used to combine the 5' of the gag-coding sequence from pGAG3 and pGAG4 with the 3' of the gag coding sequence of pR2, to generate pGAG10 and pGAG9, respectively. pGAG10 places HIV sequence from nucleotides 227 to 2037 under transcriptional control of the tryptophan promoter. pGAG9 places HIV sequence from nucleotides 273 to 2037 under tryptophan promoter control (see Figure 1). *E. coli* translational control signals for translation initiation at the gag ATG at nucleotide 334 are provided by the HIV sequences immediately 5' of the gag ATG. These translation initiation signals were predicted to be functional in *E. coli* on inspection of the HIV sequences.

Plasmid pGAG55 (Figure 2) is constructed by ligation of the following three DNA segments: 1) the large P_L-containing BamHI to NcoI fragment from pBF106; 2) the HIV gag-containing ClaI to BamHI DNA segment from pGAG9 or pGAG10; 3) a NcoI to ClaI oligonucleotide of the following sequence: 5'- C ATG GGT GCT AGA GCG TCA GTA TTA AGC GGG GGA GAA TTA GAT-3' 3'- CCA CGC TCT CGC AGT CAT AAT TCG CCC CCT CTT AAT CTA GC-5'. The NcoI to ClaI oligonucleotide contains sequence information coding for the first 14 N-terminal amino acids of gag. Plasmid pGAG55 thereby contains the complete HIV_{BH10} gag coding sequence fused to the synthetic *E. coli* translation initiation signal in plasmid pBF106, as described previously (see p. 14). Plasmid pGAG55 is also referred to as pBF128. A unique NcoI site has been engineered at the translation start ATG of gag without altering the gag coding sequence (see Figure 2). The complete gag coding sequence can now be conveniently moved from GAG55 as a NcoI to BamHI DNA fragment cassette to other expression vectors. For example, the gag coding sequence can be precisely fused to signals directing the expression of the full length gag precursor in mammalian cells. Importantly, HIV sequences 5' to the gag coding sequence have been eliminated in the NcoI to BamHI DNA cassette. HIV sequences immediately 5' to the gag coding sequence, including a splice donor signal, are very likely to limit expression of gag in mammalian cells.

Plasmid GAG17, which specifies the expres-

sion in *E. coli* of the structure, the p17^{gag} protein, was constructed by ligating the following three DNA sequences: 1) a *Sal*I-derived, approximately 2962 base pair *Sal*I-PvuII DNA fragment from pGAG55 (see Figure 1); 2) a PstI to PvuII fragment from pGAG55 (approximately 3068 base pair) containing pBR322-derived DNA, lambda-derived DNA, and the 5' portion of HIV *gag* (see Figure 2); 3) a chemically synthesized PvuII to *Sal*I DNA segment of the HIV *gag* gene (only 1 DNA strand is shown):

5'- CT GAC ACA GGA ... GAG GTC
AGC CAA AAT TAC T ... *Sal*I to *Sal*I
fragment extends the *gag* coding sequence from the PvuII site at nucleotide ... (Figure 1) and introduces of TAA stop codon ... Tyr codon (TAC) at the p17^{gag}-p24^{gag} junction (Figure 1). The 3' G residue of the indicated ... is the first nucleotide of a *Sal*I site. In ... a coding sequence corresponding ... nucleotides 334 to 729 of HIV₈₉₁₀ ... under transcriptional and translation ... derived from pBF106. pBF106 is ...

EXAMPLE 3

Reactivity of *E. coli*-expressed GAG55 with HIV-Specific Antibodies

As shown in Figure 1, GAG55 can be efficiently expressed at high levels in *E. coli*, using the appropriate plasmid vector and expression system. Analysis of total *E. coli* proteins by SDS-PAGE and Coomassie blue staining indicates that GAG55 accumulates to a level of approximately 1% of total *E. coli* protein. This high level of expression of GAG55 facilitates purification of the protein. GAG17 similarly accumulates to a high level in *E. coli*.

Monoclonal antibodies to HIV-expressed p17^{gag} (BT2, Biotec Research Labs) and p24^{gag} (BT3, Biotec Research Labs) were found to react specifically with GAG55 when tested in immunoblot and ELISA methods described previously. In addition, a rabbit antiserum specific for HIV p24^{gag} (obtained from D. R. Brown, Experimental Station) was immunoreactive to expressed GAG55. Furthermore, the reactivity of GAG55 was shown to be immunoreactive to many human AIDS patient sera in an immunoblot assay (for example, see Figure 2). *E. coli*-expressed GAG17 is immunoreactive to p17^{gag}-

specific monoclonal antibody BT2 and many human AIDS patient sera. These results confirm that viral antigenic sites reside within *E. coli*-expressed GAG55 and GAG17.

EXAMPLE 3

Use of *E. coli*-expressed GAG55 as an Immunogen

Another advantage of the antigenic structure of *E. coli*-expressed GAG55 is the ability of the peptide to illicit HIV-specific antibodies in animals. Purified GAG55 was injected into rabbits and goats and the resulting antisera were assayed by immunoblot and ELISA, according to the methods described previously. Goat and rabbit antisera to *E. coli*-expressed GAG55 reacted with the expected HIV-expressed *gag*-derived proteins p55^{gag}, p24^{gag}, and p17^{gag} on viral immunoblot. Antisera to GAG55 also reacted with an HIV-ELISA (Du Pont).

The titer of the antisera to *E. coli*-expressed GAG55 was found to be equal to or greater than antisera to *gag* proteins purified from HIV in specifically binding to the HIV-expressed *gag* proteins. This result demonstrates that *E. coli*-expressed GAG55 contains the major native viral epitopes and that antibodies to GAG55 are sensitive reagents for detecting viral proteins.

EXAMPLE 4

Detection of Antibodies to HIV *gag* in Human Sera

The *E. coli*-expressed *gag*-derived protein GAG55 is comprised of protein sequences covering the complete *gag* open reading frame and should, therefore, be very similar antigenically to P55^{gag} expressed by HIV in infected mammalian cells. In order to compare the ability of GAG55 and HIV-expressed *gag* antigens to be recognized by human antibodies in HIV-seropositive sera, experiments involving competition between recombinant and viral *gag* antigens for reaction with serum antibodies were carried out. Using a competition immunoblot technique, six HIV-seropositive human sera were tested for reactivity with GAG55 antigen as described below. Sera were pre-absorbed with GAG55-containing *E. coli* extract and, in parallel, with a control *E. coli* extract. These absorptions were followed by analysis of the sera using HIV

antigen strips (Biotech Research Labs). With each HIV positive serum tested, a recombinant gag polypeptide was found to substantially, if not completely, out-compete reactivity of all the gag-related immunoblot bands (see Figure 5). The specificity of this competition effect for gag was shown by the lack of competition of env or pol bands and the lack of effect of control E. coli extracts on any HIV immunoblot band. The present results demonstrate that most antibodies in HIV-positive sera reacting with HIV gag immunoblot bands also react with GAG55 antigen. Thus, E. coli-expressed GAG55 is clearly a useful reagent for the detection of HIV gag-specific antibodies and, therefore, can be a useful component of HIV diagnostic and blood screening procedures. These procedures may involve, for example, immunoblot or ELISA methods.

EXAMPLE 6

Purification of gag-specific antibodies in human HIV-seropositive serum by affinity chromatography using E. coli-expressed GAG

E. coli-expressed GAG55 was demonstrated using immunoblot assays to bind to essentially all of the gag-specific antibodies in many human HIV-seropositive sera (Example 5, Figure 5). Therefore, we prepared an affinity column for the preferential isolation of HIV gag-specific antibodies. GAG55 was partially purified as described above and covalently coupled to a Rane gel support (Rane Chemical, Rockford, IL) as modified by the supplier. Human serum was loaded to a column containing the GAG55 affinity and the column was washed extensively using phosphate buffered saline. Human antibodies were eluted by washing the eluted fractions were pH 8.0. Immunoblot analysis of HIV antigens (Biotech Research Labs, Rockville, MD) using the GAG55-purified human antibodies revealed selective immunoreactivity of these antibodies with gag-related antigens. These results demonstrate the utility of E. coli-expressed gag protein for the preparation of gag-specific antibodies.

HIV gag-specific antibodies are useful as research reagents for the identification and mapping of gag-specific epitopes and for the study of the host immune response to HIV. Furthermore, it may be possible to identify gag-specific antibodies that can neutralize or block HIV infection. The utility of such HIV-neutralizing antibodies as diagnostic reagents will depend on the availability and their purification. As discussed above, full

length gag protein, such as E. coli-expressed GAG55, that represents essentially all of the gag-specific epitopes may be the preferred affinity reagent for the identification and purification of gag-specific antibodies. Subfragments of gag protein, as previously disclosed by others, can only represent a subset of the total set of gag-specific epitopes.

EXAMPLE 6

Identification of HIV gag-reactive, env- and pol-nonreactive Sera

ELISAs utilizing as antigen purified HIV grown in tissue culture (HIV-ELISA) have proved valuable for screening blood for evidence of exposure to HIV. Many times sera will repeatedly test positive on HIV-ELISA but will not show clear evidence of HIV exposure when subsequently examined by HIV-immunoblot. During screening of blood donors, we and others [Courouce et al., *Lancet* II:921-922 (1986); Biberfeld et al., *Lancet* II:289-298 (1986)] often detect reactivity in HIV immunoblots at positions of gag antigens p15, p17, p24, or p55 unaccompanied by detectable immunoreactivity with env. Examples of such HIV gag-reactive, env-non-reactive sera are shown in Figure 5. The specificity of these human antibodies for HIV gag was demonstrated by competition binding with GAG55. Thus, preincubation of the sera with GAG55 specifically blocked binding of the antibody to the gag proteins on the HIV-immunoblot.

More evidence demonstrating the occurrence of anti-gag antibodies and quantitative data on their relative levels in different populations was obtained using purified recombinant antigens in ELISA format as shown in Figure 6. The HIV-ELISA-reactive blood donors are distinct from the normal group in having elevated gag ELISA titres, as does the HIV-positive group. In addition, the GAG55-ELISA values for the HIV-ELISA-reactive blood donors are in the same range as several of the HIV-positive sera. Analysis of these samples with the ENV9-ELISA revealed that the absorbances of the HIV-ELISA reactive donor group were distinctly lower than the HIV-positive group and similar to those of normal sera.

For a high percentage of the more than 200 HIV-ELISA positive sera that we have examined, there is unequivocal evidence of antibodies that are immunoreactive with HIV gag antigens, unaccompanied by evidence of HIV env reactivity. The methods demonstrated utilizing both GAG55-ELISA

and ENV9-ELISA should prove to be a routine blood screening and diagnostic assay.

The ability to identify such gag-seropositive and env-seronegative human samples is of considerable importance. For example, several studies, including that of A. S. [unintelligible] presented at the National Institutes of Health AIDS Development Conference on "The Role of Routine HTLV-III Antibody Testing in the Blood Bank" (July 7-9, 1986), have indicated that individuals who initially display immuno-reactivity only to gag specific bands may subsequently display definite HIV seropositivity if env reactivity. It is also possible that HIV gag-reactive, env-nonreactive sera may be due to other human retroviruses, such as HTLV-I, HTLV-II or HIV-II, and possibly to non-HIV retroviruses. The significance to the transfusion recipient of such HIV gag-reactive and env-non-reactive sera remains to be determined, and further studies are essential. As a first step, blood from donors with these sera should not be transfused. This example illustrates both the importance of methods for the detection of HIV gag proteins and the utility of GAG55 ELISA.

EXAMPLE 7

Analysis of Human Antisera to gag and env Expressed env- and gag-specific Antigens

Although the HIV ELISA is a powerful and sensitive tool for screening for the presence of HIV-specific antibodies, utilizing total virus antigen information concerning which components of the virus are immunogenic in sera. Immunoblot or immunoprecipitation can be used to identify immunoreactivity to specific viral components. To date, the detection of HIV-seropositivity is based on reactivity with both gag- and env-specific antigens.

As an alternative to the immunoprecipitation method for determining relative levels of antibodies to gag and env proteins, we have developed gag- and env-specific ELISAs, using recombinant GAG55 and ENV9 as antigens. ENV9 corresponds to the amino acids corresponding to the C-terminus of the HIV_{ENV} protein (p17 and 240 amino acids of the N-terminal region of the main chain). Methods for detection of relative levels of antibodies to these antigenic components of HIV are described in Example 6.

encoded proteins, provides information that may be useful for the diagnosis and prognosis of HIV-infected individuals. The ELISA format provides considerable advantages compared to immunoblot and immunoprecipitation methods. The ELISA procedure is sensitive, quantitative, rapid, non-radioactive, and relatively inexpensive and simple to run.

Recently several ELISA assays have been developed which utilize as antigen proteins expressed in *E. coli* to detect antibodies to the AIDS virus [Dowbenko, et al., *Proc. Natl. Acad. Sci.* 82: 7748-7752 (1985); Steimer, et al., *Virology* 150:283-290 (1986); Cabradilla et al., *Biotechnology* 4: 128-133 (1986); Chang et al., *Biotechnology* 3:905-909 (1985); Shoeman et al., *Anal. Biochem.* 161:370-379 (1987)]. Previous recombinant gag-derived ELISAs which have been described only contain subregions of gag, such as p24^{gag}, and do not contain sequences corresponding to the entire gag coding sequence. Since the GAG55 represents the complete gag coding sequence, the GAG55-ELISA may detect HIV gag-specific antibodies in human sera that are not detected by ELISAs employing subregions of gag. In particular, GAG55 is preferred to other reported recombinant gag antigens because the complete p17^{gag} region is represented. In addition, GAG55 and GAG17 do not contain any non-HIV or non-HIV gag amino acid sequences. These antigens are preferred for antibody detection because of the specificity for gag and of the reduced probability of detecting false-positive, non-HIV immunoreactivity.

The HIV gag-specific and env-specific antibody levels were quantitated in human sera from individuals of known clinical status, using both the GAG55-ELISA and ENV9-ELISA. Sera categorized as being from AIDS patients, HIV-seropositive (non-AIDS), and HIV-seronegative individuals were obtained from Duke University. These sera were analyzed using the GAG55 and ENV9 ELISAs. The results of these assays are shown in Figure 7.

As shown in Figure 7, the gag-specific apparent antibody levels in many of the AIDS patient sera are reduced compared to the gag-specific apparent antibody levels in non-AIDS, HIV-seropositive sera. This effect is selective, since the HIV env-specific antibody levels do not distinguish the AIDS and non-AIDS seropositive groups. The GAG55-ELISA alone is obviously limited as an HIV screening assay since many of the characterized HIV positive sera were not detected as positive (see also Example 6, Figure 6). This does not, however, preclude the use of a gag-specific ELISA as an important component of diagnostic assays. For example, many sera exhibit HIV gag immunoreactivity without detectable env-specific immunoreactivity (Example 6). Moreover, quantitation of the relative antibody levels specific to gag and

env and other HIV antigenic components may be useful for monitoring HIV-associated disease. The results presented indicate that it may be possible to develop a correlation between the serum reactivity on the GAG55 and ENV ELISA assays and the stage of disease. The present data provide evidence that quantitation by ELISA of the relative HIV env-specific and gag-specific antibody levels in human sera is useful for both screening and diagnostic analysis of human sera.

Claims

1. A recombinant plasmid capable of high level expression in *E. coli* of a nonfused peptide corresponding to the complete HIV coding sequence and exhibiting the antigenicity of the complete HIV p55^{gag} protein, consisting essentially of plasmid pBR322 and DNA coding for amino acids 1 to 512 of HIV protein p55^{gag}, with a translation start signal and an ATG/Met translation start codon downstream from a promoter selected from the group consisting of the *E. coli* lac operon promoter, the *E. coli* tryptophan operon promoter, and the phage lambda P_L promoter.

2. A recombinant plasmid of Claim 1 having incorporated therein a translation start signal which is an HIV sequence immediately 5' of the gag ATG/Met translation start codon or an oligonucleotide linker as defined in Example 1.

3. A recombinant plasmid of Claim 1 wherein the gag ATG/Met translation start codon is part of an NcoI restriction endonuclease site.

4. An *E. coli* cell transformed with a plasmid of anyone of claims 1 to 3.

5. An HIV gag-encoded peptide expressed by a cell of Claim 4.

6. A recombinant plasmid capable of high level expression in *E. coli* of a nonfused peptide corresponding to the complete HIV p17^{gag} coding sequence and exhibiting the antigenicity of the complete HIV p17^{gag} protein, consisting essentially of plasmid pBR322 and DNA coding for amino acids 1 to 132 of HIV protein p55^{gag}, with a translation start signal and an ATG/Met translation start codon downstream from a promoter selected from the group consisting of the *E. coli* lac operon promoter, the *E. coli* tryptophan operon promoter, and the phage lambda P_L promoter.

7. A recombinant plasmid of Claim 6 having incorporated therein a translation start signal which is an HIV sequence immediately 5' of the gag ATG/Met translation start codon or an oligonucleotide linker as defined in Example 1.

8. A recombinant plasmid of Claim 6 wherein the gag ATG/Met translation start codon is part of an NcoI restriction endonuclease site.

9. An *E. coli* cell transformed with a plasmid of anyone of claims 6 to 8.

10. An HIV gag-encoded peptide expressed by a cell of Claim 9.

11. An *E. coli*-expressed recombinant nonfused peptide corresponding to the complete HIV p55^{gag} coding sequence and exhibiting the antigenicity of the complete HIV p55^{gag} protein.

12. An *E. coli*-expressed recombinant nonfused peptide according to Claim 11 which is encoded by nucleotides numbered 334 to 1869 of the HIV genome, as shown in Figure 1.

13. An *E. coli*-expressed recombinant nonfused peptide corresponding to the complete p17 sub-region of the HIV p55^{gag} coding sequence, and exhibiting the antigenicity of the complete HIV p17^{gag} protein.

14. An *E. coli*-expressed recombinant nonfused peptide according to Claim 13 which is encoded by nucleotides numbered 334 to 729 of the HIV genome, as shown in Figure 1.

15. In a diagnostic kit used for detecting antibodies to HIV in a biological sample wherein the sample is contacted with a peptide and immunoreactivity is detected, the improvement comprising employing a peptide of anyone of claims 5, 10, 11, and 13.

16. In a process for detecting antibodies to HIV in a biological sample, comprising contacting said sample with a peptide which is immunoreactive with said antibodies and detecting immunoreactivity, the improvement comprising employing a peptide of anyone of claims 5, 10, 11, and 13.

17. A vaccine protective against HIV comprising an effective protective amount of a peptide of anyone of claims 5, 10, 11, and 13 in a pharmaceutically acceptable carrier.

F I G. 1A

228 255
 GAA AGC GGA A B C GAG AGC TCT CTC GAC GCA GGA CTC GGC TTG CTG AAG
 Glu Ser Gly Lys G. Arg Ser Ser Leu Asp Ala Gly Leu Gly Leu Leu Lys

282 309
 CGC GCA CGG C A G G G GGC GAC TGG TGA GTA CGC CAA AAA TTT TGA
 Arg Ala Arg Glu C G Gly Gly Asp Trp . Val Arg Gln Lys Phe .

336 363
 CTA GCG GAG C A G AG GAG ATG GGT GCG AGA GCG TCA GTA TTA AGC GGG
 Leu Ala Glu Glu Arg. Arg. Glu MET Gly Ala Arg Ala Ser Val Leu Ser Gly

390 417
 GGA GAA TTA C A G G A AAA ATT CGG TTA AGG CCA GGG GGA AAG AAA AAA
 Gly Glu Leu Arg Lys Lys Ile Arg Leu Arg Pro Gly Gly Lys Lys Lys

444 471
 TAT AAA TTA A A G GT. TGG GCA AGC AGG GAG CTA GAA CGA TTC GCA GTT
 Tyr Lys Leu Lys Val. Trp Ala Ser Arg Glu Leu Glu Arg Phe Ala Val

498 525
 AAT CCT GGC C A G TCA GAA GGC TGT AGA CAA ATA CTG GGA CAG CTA
 Asn Pro Gly Lys Ser Glu Gly Cys Arg Gln Ile Leu Gly Gln Leu

552 579
 CAA CCA TCC C A G GG TCA GAA GAA CTT AGA TCA TTA TAT AAT ACA GTA
 Gln Pro Ser Leu Gl. Ser Glu Glu Leu Arg Ser Leu Tyr Asn Thr Val

606 633
 GCA ACC CTC T A G CAA AGG ATA GAG ATA AAA GAC ACC AAG GAA GCT
 Ala Thr Leu Thr Gln Arg Ile Glu Ile Lys Asp Thr Lys Glu Ala

660 687
 TTA GAC AAG C A G CAA AAC AAA AGT AAG AAA AAA GCA CAG CAA GCA
 Leu Asp Lys Thr Gln Gln Asn Lys Ser Lys Lys Lys Ala Gln Gln Ala

714 741
 GCA GCT GAC A A G AGT CAG GTC AGC CAA AAT TAC CCT ATA GTG CAG
 Ala Ala Asp Thr Ser Gln Val Ser Gln Asn Tyr Pro Ile Val Gln

768 795
 AAC ATC CAG C A G CAT CAG GCC ATA TCA CCT AGA ACT TTA AAT GCA
 Asn Ile Gln Lys G. His Gln Ala Ile Ser Pro Arg Thr Leu Asn Ala

822 849
 TGG GTA AAA C A G AAG GCT TTC AGC CCA GAA GTA ATA CCC ATG TTT
 Trp Val Lys Val Lys Ala Phe Ser Pro Glu Val Ile Pro MET Phe

F I G. 1 B

876 903
TCA GCA TTA TCA GAA GGA ACC ACC CCA CAA GAT TTA AAC ACC ATG CTA AAC ACA
Ser Ala Leu Ser Glu Gly Ala Thr Pro Gln Asp Leu Asn Thr MET Leu Asn Thr

930 957
GTG GGG GGA CAT CC GCA GGC ATG CAA ATG TTA AAA GAG ACC ATC AAT GAG GAA
Val Gly Gly His Gl Ala Ala MET Gln MET Leu Lys Glu Thr Ile Asn Glu Glu

984 1011
GCT GCA GAA TGG GAT AGA CTA CAT CCA GTG CAT GCA GGG CCT ATT GCA CCA GGC
Ala Ala Glu Trp Asp Arg Val His Pro Val His Ala Gly Pro Ile Ala Pro Gly

1038 1065
CAG ATG AGA GAA CAA AGG GAA AGT GAC ATA GCA GGA ACT ACT AGT ACC CTT CAG
Gln MET Arg Glu Pro Arg Gly Ser Asp Ile Ala Gly Thr Thr Ser Thr Leu Gln

1092 1119
GAA CAA ATA GCA TCG ATG CAA AAT AAT CCA CCT ATC CCA GTA GGA GAA ATT TAT
Glu Gln Ile Gly Trp MET Thr Asn Asn Pro Pro Ile Pro Val Gly Glu Ile Tyr

1146 1173
AAA AGA TGG ATA CCG CTG CAA TTA AAT AAA ATA GTA AGA ATG TAT AGC CCT ACC
Lys Arg Trp Ile Thr Leu Gly Leu Asn Lys Ile Val Arg MET Tyr Ser Pro Thr

1200 1227
AGC ATT CTG GAG AAT AGA CAA GGA CCA AAA GAA CCT TTT AGA GAC TAT GTA GAC
Ser Ile Leu Asp Ile Arg Gln Gly Pro Lys Glu Pro Phe Arg Asp Tyr Val Asp

1254 1281
CGG TTC TAT AAA CTA CAA GCC GAG CAA GCT TCA CAG GAG GTA AAA AAT TGG
Arg Phe Tyr Lys Thr Leu Arg Ala Glu Gln Ala Ser Gln Glu Val Lys Asn Trp

1308 1335
ATG ACA GAA AAT CCG TTG CCG CAA AAT GCG AAC CCA GAT TGT AAG ACT ATT TTA
MET Thr Glu Thr Leu Leu Val Gln Asn Ala Asn Pro Asp Cys Lys Thr Ile Leu

1362 1389
AAA GCA TTG GAA CCG GCT ACA CTA GAA GAA ATG ATG ACA GCA TGT CAG GGA
Lys Ala Leu Gly Pro Ala Ala Thr Leu Glu Glu MET MET Thr Ala Cys Gln Gly

1416 1443
GTA GGA GGA CCG GCG CAT AAG GCA AGA GTT TTG GCT GAA GCA ATG AGC CAA GTA
Val Gly Gly Pro Gly His Lys Ala Arg Val Leu Ala Glu Ala MET Ser Gln Val

24

1470 1497
ACA AAT ACA GAA CTA ATG ATG CAG AGA GGC AAT TTT AGG AAC CAA AGA AAG
Thr Asn Thr Ala Thr Ile MET MET Gln Arg Gly Asn Phe Arg Asn Gln Arg Lys

1524 1551
ATG GTT AAG TCA TCA AAT TCT GGC AAA GAA GGG CAC ACA GCC AGA AAT TGC AGG
MET Val Lys Cys Phe Asn Cys Gly Lys Glu Gly His Thr Ala Arg Asn Cys Arg

F I G. 1 C

1578 1605
GCC CCT AGG AAA AAG GGC TGT TGG AAA TGT GGA AAG GAA GGA CAC CAA ATG AAA
Ala Pro Arg Lys Lys Gly Cys Trp Lys Cys Gly Lys Glu Gly His Gln MET Lys

1632 1659
GAT TGT ACT GAG AGA CAG GCT AAT TTT TTA GGG AAG ATC TGG CCT TCC TAC AAG
Asp Cys Thr Glu Arg Gln Ala Asn Phe Leu Gly Lys Ile Trp Pro Ser Tyr Lys

1686 1713
GGA AGG CCA GGG AAT TTT CTT CAG AGC AGA CCA GAG CCA ACA GCC CCA CCA TTT
Gly Arg Pro Gly Asn Phe Leu Gln Ser Arg Pro Glu Pro Thr Ala Pro Pro Phe

1740 1767
CTT CAG AGC AGA CCA GAG CCA ACA GCC CCA CCA GAA GAG AGC TTC AAG TCT GGG
Leu Gln Ser Arg Phe Glu Pro Thr Ala Pro Pro Glu Glu Ser Phe Arg Ser Gly

1794 1821
GTA GAG ACA ACA ACT CCC CCT CAG AAG CAG GAG CCG ATA GAC AAG GAA CTG TAT
Val Glu Thr Thr Thr Pro Pro Gln Lys Gln Glu Pro Ile Asp Lys Glu Leu Tyr

1848 1875
CCT TTA ACT TCC CTT AGA TCA CTC TTT GGC AAC GAC CCC TCG TCA CAA TAA AGA
Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp Pro Ser Ser Gln Arg

1902 1929
TAG GGG GGC AAC TAA AGG AAG CTC TAT TAG ATA CAG GAG CAG ATG ATA CAG TAT
Gly Gly Asn Arg Lys Leu Tyr Ile Gln Glu Gln MET Ile Gln Tyr

1956 1983
TAG AAG AAA TGA CTT TGC CAG GAA GAT GGA AAC CAA AAA TGA TAG GGG GAA TTG
Lys Lys Val Lys Gln Glu Asp Gly Asn Gln Lys Gly Glu Leu

2010
GAG GTT TTA TCA AAT TA
Glu Val Leu Ser Lys

FIG. 2

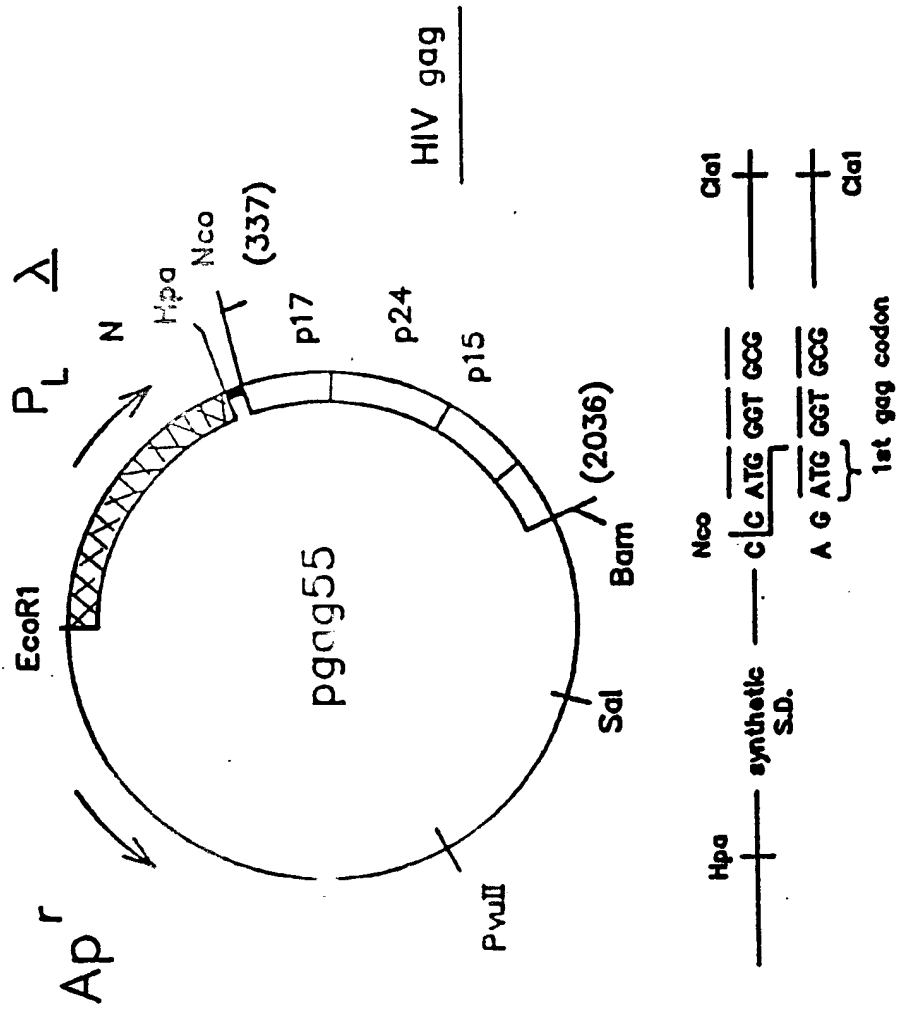
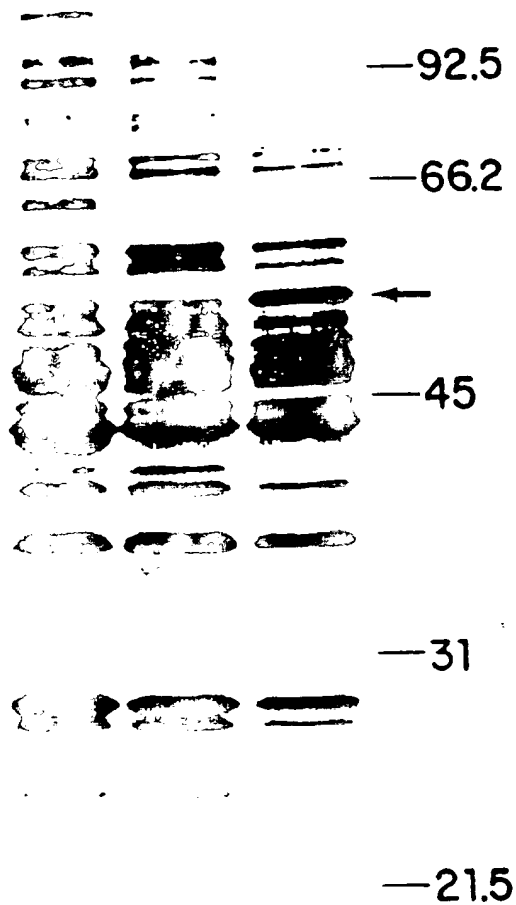


FIG. 3

1 2 3



F I G. 4

1 2 3

—97.4

—68

—43

—25.7

—18.4

F I G. 5

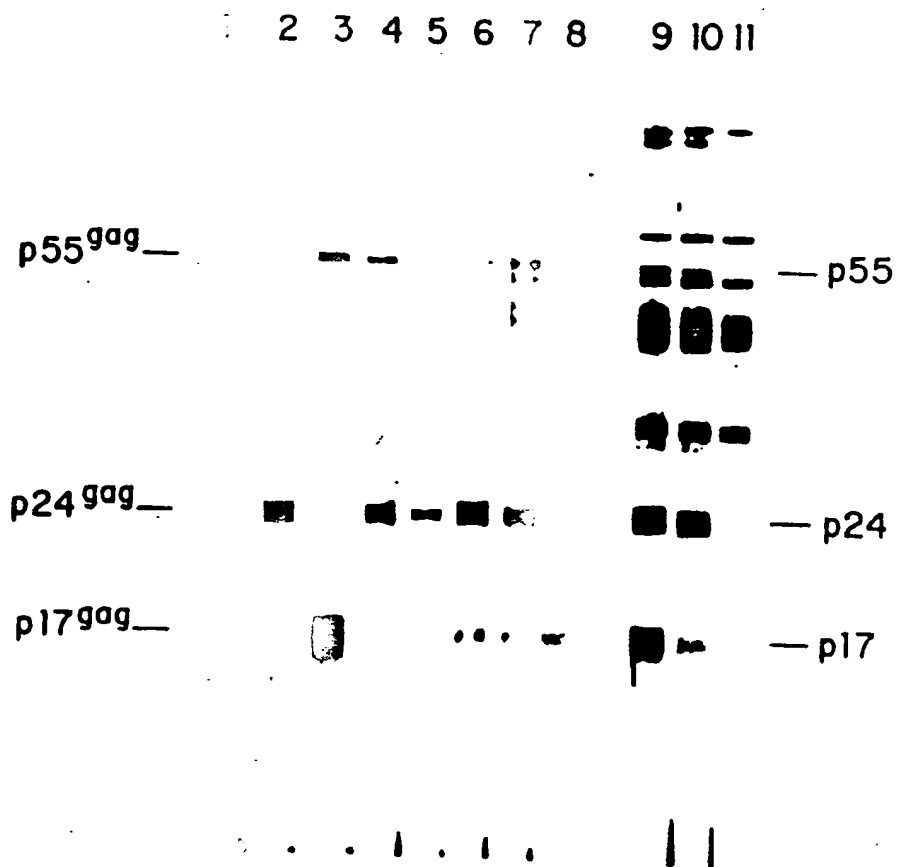


FIG. 6A

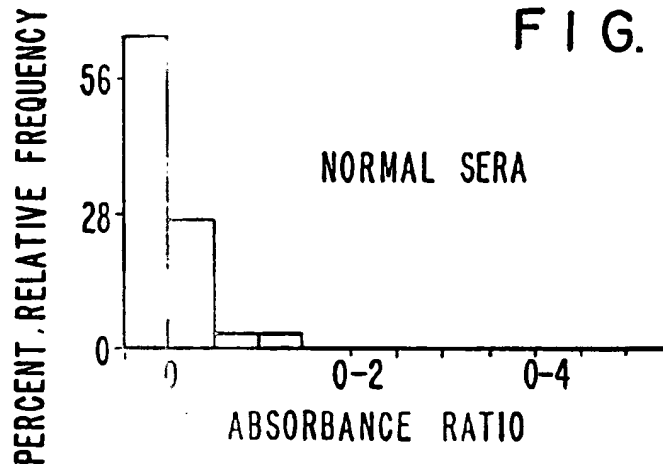


FIG. 6B

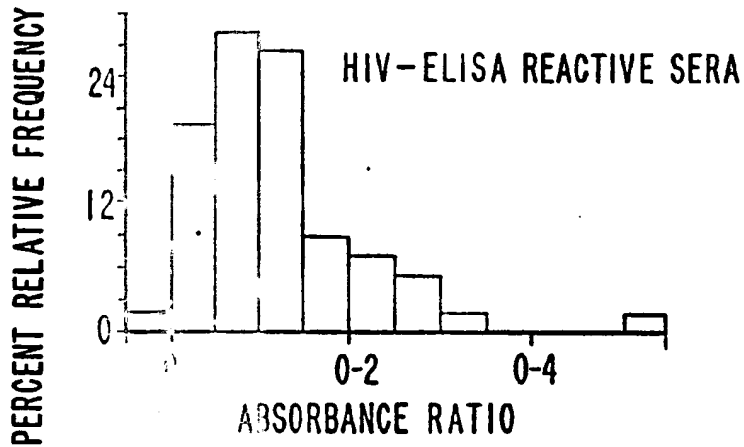


FIG. 6C

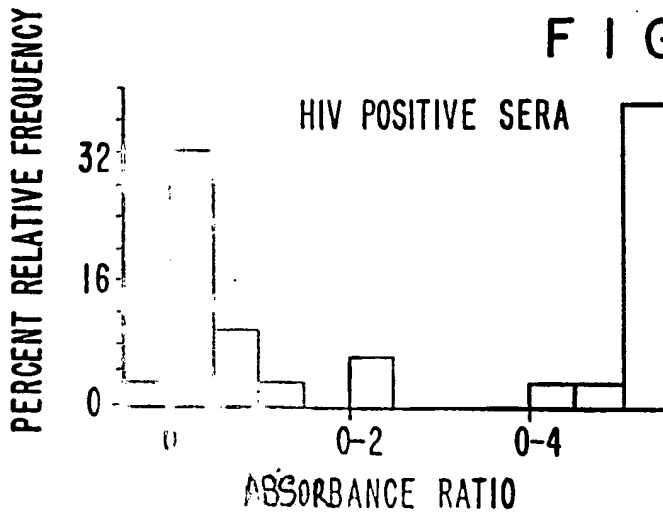
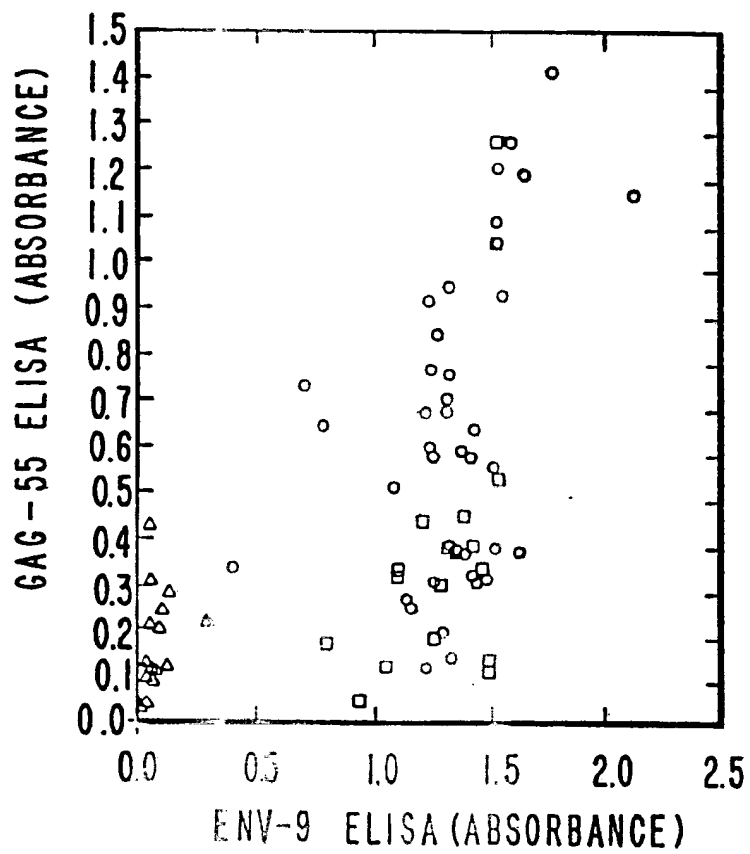


FIG. 7



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